

#3

## SEQUENCE LISTING

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Sangamo BioSciences, Inc.

<120> Regulation of Angiogenesis With Zinc  
Finger Proteins

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1 5

<210> 181  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> finger

<400> 181  
Gln Ser Ser Asp Leu Thr Arg  
1 5

<210> 182  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> target

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<220>
<223> target
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<210> 184
<211> 9
<212> DNA
<213> Artificial Sequence

<220>
<223> target
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<210> 185
<211> 9
<212> DNA
<213> Artificial Sequence
<220>
<223> target
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<210> 186
<211> 7
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<213> Artificial Sequence

<220>
<223> recognition helix
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<210> 187
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> recognition helix
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<400> 187  
 Gln Ser Gly Asp Leu Thr Arg  
 1 5

<210> 188  
 <211> 7  
 <212> PRT  
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<220>  
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<400> 188  
 Glu Arg Gly Asp Leu Thr Arg  
 1 5

<210> 189  
 <211> 7  
 <212> PRT  
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<220>  
 <223> recognition helix

<400> 189  
 Arg Ser Asp His Leu Ala Arg  
 1 5

<210> 190  
 <211> 7  
 <212> PRT  
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<220>  
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<400> 190  
 Arg Ser Asp Asn Leu Ala Arg  
 1 5

<210> 191  
 <211> 7  
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<220>  
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<400> 191  
 Gln Ser Ser His Leu Ala Arg  
 1 5

<210> 192  
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<220>  
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<400> 192  
 Arg Ser Asp Glu Leu Thr Arg  
 1 5

<210> 193  
 <211> 7  
 <212> PRT  
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<220>  
 <223> recognition helix

<400> 193  
 Arg Ser Asp Glu Leu Gln Arg  
 1 5

<210> 194  
 <211> 7  
 <212> PRT  
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<220>  
 <223> recognition helix

<400> 194  
 Arg Ser Asp Asn Leu Ala Arg  
 1 5

<210> 195  
 <211> 7  
 <212> PRT  
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<220>  
 <223> recognition helix

<400> 195  
 Arg Ser Asp His Leu Ala Arg  
 1 5

<210> 196  
 <211> 7  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> recognition helix

<400> 196  
 Asp Arg Ser Asn Leu Thr Arg  
 1 5

<210> 197  
 <211> 7  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> recognition helix

<400> 197  
 Arg Ser Asp Ala Leu Thr Gln  
 1 5

$\langle 400 \rangle$  202  
 Arg Ser Asp His Leu Thr Arg  
 1 5





10

<213> Artificial Sequence

class of zinc finger proteins (ZFP)

<223> Xaa = any amino acid

may be present or absent

1                      5  
Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His  
                20                      25

<213> Artificial Sequence

<223> t'arget

ggcgtagac

<213> Artificial Sequence

<223> target

ggcgacgta

<213> Artificial Sequence

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<220>
<223> peptide linker

<400> 211
Thr Gly Glu Lys Pro
1          5

<210> 212
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> peptide linker

<400> 212
Gly Gly Gly Gly Ser
1          5

<210> 213
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> peptide linker

<400> 213
Gly Gly Arg Arg Gly Gly Gly Ser
1          5

<210> 214
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> peptide linker

<400> 214
Leu Arg Gln Arg Asp Gly Glu Arg Pro
1          5

<210> 215
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> peptide linker

<400> 215
Leu Arg Gln Lys Asp Gly Gly Gly Ser Glu Arg Pro
1          5          10

<210> 216
<211> 16
<212> PRT
<213> Artificial Sequence

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<221> modified_base
<222> (1)...(1)
<223> n = c modified by aminofluorescein (FAM)
```

```

<221> modified_base
<222> (25)...(25)
<223> n = a modified by tetramethylrhodamine (TAMRA)

<400> 228
nagtagctgc gctgatagac atccn
25

<210> 229
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> GAPDH forward primer

<400> 229
ccatgttcgt catgggtgtg a
21

<210> 230
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> GAPDH reverse primer

<400> 230
catggactgt ggcatgagt
20

<210> 231
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> GAPDH Taqman probe

<221> modified_base
<222> (1)...(1)
<223> n = t modified by aminofluorescein (FAM)

<221> modified_base
<222> (24)...(24)
<223> n = a modified by tetramethylrhodamine (TAMRA)

<400> 231
ncctgcacca ccaactgctt agcn
24

<210> 232
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> VP16-FLAG forward primer

<400> 232
catgacgatt tcgatctgga
20

<210> 233
<211> 22

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<212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> VP16-FLAG reverse primer  
  
 <400> 233  
 ctacttggtca tcgtcgtcct tg 22  
  
 <210> 234  
 <211> 26  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> VP16-FLAG Tagman probe  
  
 <221> modified\_base  
 <222> (1)...(1)  
 <223> n = a modified by aminofluorescein (FAM)  
  
 <221> modified\_base  
 <222> (26)...(26)  
 <223> n = a modified by tetramethylrhodamine (TAMRA)  
  
 <400> 234  
 ntcggtaaac atctgctcaa actcgn 26  


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 <210> 235  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> RT-PCR primer  
  
 <400> 235  
 atgaactttc tgctgtcttg ggtgcatt 28  
  
 <210> 236  
 <211> 22  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> RT-PCR primer  
  
 <400> 236  
 tcaccgcctc ggcttgtcac at 22  
  
 <210> 237  
 <211> 18  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> murine VEGF target  
  
 <400> 237  
 tgagcggcgg cagcggag 18



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<210> 243
<211> 18
<212> PRT
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<212> PRT

<213> Artificial Sequence

<220>

<223> finger

<400> 248

Thr Ser Gly His Leu Ala Arg  
1 5

<210> 249

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> finger

<400> 249

Thr Ser Gly His Leu Arg Arg  
1 5

<210> 250

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> finger

<400> 250

Thr Ala Gly His Leu Val Arg  
1 5

<210> 251

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> finger

<400> 251

Thr Thr Gly His Leu Val Arg  
1 5

<210> 252

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> finger

<400> 252

Thr Lys Asp His Leu Val Arg  
1 5